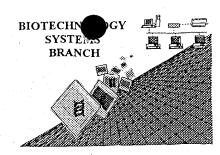
North

## RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09/823, 257
Source:	OIPE
Date Processed by STIC:	4-17-01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: <u>patin21help@uspto.gov</u> or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: <u>patin3help@uspto.gov</u> or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

## Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

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RAW SEQUENCE LISTING
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PATENT APPLICATION: US/09/823,257

DATE: 04/17/2001 TIME: 10:50:13

Input Set : A:\P0715seqUS.txt

Output Set: N:\CRF3\04172001\1823257.raw

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      5 <120> TITLE OF INVENTION: High Throughput Methods for Haplotyping
      7 <130> FILE REFERENCE: P0715/7003 (HCL)
C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/823,257
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     9 <150> PRIOR APPLICATION NUMBER: US 60/194,425
     10 <151> PRIOR FILING DATE: 2000-04-04
     12 <160> NUMBER OF SEQ ID NOS: 24
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     14 <170> SOFTWARE: PatentIn version 3.0
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    21 <220> FEATURE:
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    70 <213> ORGANISM: Homo sapiens
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    note V
                                                                                   Oligonucleotide.
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The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/823,257

DATE: 04/17/2001 TIME: 10:50:13

Input Set : A:\P0715seqUS.txt

Output Set: N:\CRF3\04172001\1823257.raw

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/823,257

DATE: 04/17/2001 TIME: 10:50:14

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RAW SEQUENCE LISTING PATENT APPLICATION: US/09/823,257

DATE: 04/17/2001 TIME: 10:50:14

Input Set : A:\P0715seqUS.txt

Output Set: N:\CRF3\04172001\1823257.raw

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RAW SEQUENCE LISTING

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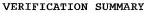
PATENT APPLICATION: US/09/823,257

Input Set : A:\P0715seqUS.txt

Output Set: N:\CRF3\04172001\1823257.raw

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PATENT APPLICATION: US/09/823,257

DATE: 04/17/2001 TIME: 10:50:15

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